```
logo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 374 SMU RS06885 WP 002262930.1          ......MKKILIVDDEKPISDIIKFNLAKEGYDTITAFDGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRKN...S    71
NTDB id 309732 DOM02 RS02700 WP 003084583.1     MEQQSDGLKVMVIDDSKTIRRTAETLLKKVGCDVITAIDGFDALAKIADTHPNIIFVDIMMPRLDGYQTCALIKNNSAFK    80
NTDB id 1047 HON27 RS03110 WP 000389061.1       MEDAFQNLKVMVIDDSKTIRRTAETLLQREGCEVITAVDGFEALSKIAEANPDIVFVDIMMPRLDGYQTCALIKNSQNYQ    80
consensus                                       **      *!****!!*!*!*******!  **!***!!!*!!**!!*!*** *!*****!*!*!*!!!**********  *

logo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 374 SMU RS06885 WP 002262930.1          HVPIIMLSAKDSEFDKVIGLEIGADDYVTKPFSNRELLARVKAHLRRTENIESAVAEENASGIPEIIIGDLQILPDAFVA   151
NTDB id 309732 DOM02 RS02700 WP 003084583.1     STPVIMLSSKDGLFDKAKGRIVGSDQYLTKPFSKEELLGAIKAHVPSFTPVDAVS........................   135
NTDB id 1047 HON27 RS03110 WP 000389061.1       NIPVIMLSSKDGLFDQAKGRVVGSDEYLTKPFSKDELLNAIRNHVSS................................   127
consensus                                        *!*!!!!*!!**!!***!***!*!*!*!!!!!**!!!*****!* *   **

logo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 374 SMU RS06885 WP 002262930.1          KKRGTEVELTHREFELLHHLATHTGQVMTREHLLETVWGYDYFGDVRTVDVTVRRLREKIEDTPSRPEYILTRRGVGYYM   231
NTDB id 309732 DOM02 RS02700 WP 003084583.1     ......................................................................................   135
NTDB id 1047 HON27 RS03110 WP 000389061.1       ......................................................................................   127
consensus

logo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 374 SMU RS06885 WP 002262930.1          KSYD   235
NTDB id 309732 DOM02 RS02700 WP 003084583.1     ....   135
NTDB id 1047 HON27 RS03110 WP 000389061.1       ....   127
consensus
```

X non conserved
X similar
X ≥ 50% conserved